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RAW SEQUENCE LISTING DATE: 11/13/2002  
PATENT APPLICATION: US/09/898,234B TIME: 15:47:18

DATE: 11/13/2002  
TIME: 15:47:18

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Output Set: N:\CRF4\11132002\I898234B.raw

3 <110> APPLICANT: Hauptmann, Rudolph  
4       Himmler, Adolph  
5       Maurer-Fogy, Ingrid  
6       Stratowa, Christian  
8 <120> TITLE OF INVENTION: TNF Receptors, TNF Binding Proteins and DNAs Coding for  
9       Them  
11 <130> FILE REFERENCE: 98-385-I  
13 <140> CURRENT APPLICATION NUMBER: 09/898,234B  
14 <141> CURRENT FILING DATE: 2001-07-03  
16 <150> PRIOR APPLICATION NUMBER: 09/525,998  
17 <151> PRIOR FILING DATE: 2000-03-15  
19 <150> PRIOR APPLICATION NUMBER: 08/383,676  
20 <151> PRIOR FILING DATE: 1995-02-01  
22 <150> PRIOR APPLICATION NUMBER: 08/153,287  
23 <151> PRIOR FILING DATE: 1993-11-17  
25 <150> PRIOR APPLICATION NUMBER: 07/821,750  
26 <151> PRIOR FILING DATE: 1992-01-02  
28 <150> PRIOR APPLICATION NUMBER: 07/511,430  
29 <151> PRIOR FILING DATE: 1990-04-20  
31 <160> NUMBER OF SEQ ID NOS: 97  
33 <170> SOFTWARE: PatentIn Ver. 2.0  
35 <210> SEQ ID NO: 1  
36 <211> LENGTH: 1368  
37 <212> TYPE: DNA  
38 <213> ORGANISM: Homo sapiens  
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42 <222> LOCATION: (1)..(1365)  
44 <220> FEATURE:  
45 <221> NAME/KEY: sig\_peptide  
46 <222> LOCATION: (1)..(87)  
48 <220> FEATURE:  
49 <221> NAME/KEY: misc\_feature  
50 <222> LOCATION: (88)..(120)  
51 <223> OTHER INFORMATION: portion of TNF-BP pro protein cleaved by  
52       extracellular proteases following secretion  
54 <220> FEATURE:  
55 <221> NAME/KEY: misc\_feature  
56 <222> LOCATION: (606)..(633)  
57 <223> OTHER INFORMATION: portion of TNF-BP pro protein cleaved by  
58       extracellular proteases following secretion  
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62	Met	Gly	Leu	Ser	Thr	Val	Pro	Asp	Leu	Leu	Leu	Pro	Leu	Val	Leu	Leu	
63	1						5			10					15		
65	gag	ctg	ttg	gtg	gga	ata	tac	ccc	tca	ggg	gtt	att	gga	ctg	gtc	cct	96
66	Glu	Leu	Leu	Val	Gly	Ile	Tyr	Pro	Ser	Gly	Val	Ile	Gly	Leu	Val	Pro	
67							20			25					30		
69	cac	cta	ggg	gac	agg	agg	aag	aga	gtt	gtg	tgt	ccc	caa	gga	aaa		144
70	His	Leu	Gly	Asp	Arg	Glu	Lys	Arg	Asp	Ser	Val	Cys	Pro	Gln	Gly	Lys	
71							35			40					45		
73	tat	atc	cac	cct	caa	aat	aat	tcg	att	tgc	tgt	acc	aag	tgc	cac	aaa	192
74	Tyr	Ile	His	Pro	Gln	Asn	Asn	Ser	Ile	Cys	Cys	Thr	Lys	Cys	His	Lys	
75							50			55					60		
77	gga	acc	tac	ttg	tac	aat	gac	tgt	cca	ggc	ccg	ggg	cag	gat	acg	gac	240
78	Gly	Thr	Tyr	Leu	Tyr	Asn	Asp	Cys	Pro	Gly	Pro	Gly	Gln	Asp	Thr	Asp	
79	65						70			75					80		
81	tgc	agg	gag	tgt	gag	agc	ggc	tcc	tcc	acc	gct	tca	gaa	aac	cac	ctc	288
82	Cys	Arg	Glu	Cys	Glu	Ser	Gly	Ser	Phe	Thr	Ala	Ser	Glu	Asn	His	Leu	
83							85			90					95		
85	aga	cac	tgc	ctc	agc	tgc	tcc	aaa	tgc	cga	aag	gaa	atg	ggt	cag	gtg	336
86	Arg	His	Cys	Leu	Ser	Cys	Ser	Lys	Cys	Arg	Lys	Glu	Met	Gly	Gln	Val	
87							100			105					110		
89	gag	atc	tct	tct	tgc	aca	gtg	gac	cg	gac	acc	gtg	tgt	ggc	tgc	agg	384
90	Glu	Ile	Ser	Ser	Cys	Thr	Val	Asp	Arg	Asp	Thr	Val	Cys	Gly	Cys	Arg	
91							115			120					125		
93	aag	aac	cag	tac	cg	cat	tat	tgg	agt	gaa	aac	ctt	tcc	cag	tgc	tcc	432
94	Lys	Asn	Gln	Tyr	Arg	His	Tyr	Trp	Ser	Gl	Asn	Leu	Phe	Gln	Cys	Phe	
95							130			135					140		
97	aat	tgc	agc	ctc	ctc	aat	ggg	acc	gtg	cac	ctc	tcc	tgc	cag	gag		480
98	Asn	Cys	Ser	Leu	Cys	Leu	Asn	Gly	Thr	Val	His	Leu	Ser	Cys	Gln	Glu	
99	145						150			155					160		
101	aaa	cag	aac	acc	gtg	tgc	acc	tgc	cat	gca	gg	ttt	ctt	cta	aga	gaa	528
102	Lys	Gln	Asn	Thr	Val	Cys	Thr	Cys	His	Ala	Gly	Phe	Phe	Leu	Arg	Glu	
103							165			170					175		
105	aac	gag	tgt	gtc	tcc	tgt	agt	aac	tgt	aag	aaa	agc	ctg	gag	tgc	acg	576
106	Asn	Glu	Cys	Val	Ser	Cys	Ser	Asn	Cys	Lys	Ser	Leu	Glu	Cys	Thr		
107							180			185					190		
109	aag	ttg	tgc	cta	ccc	cag	att	gag	aat	gtt	aag	ggc	act	gag	gac	tca	624
110	Lys	Leu	Cys	Leu	Pro	Gln	Ile	Glu	Asn	Val	Lys	Gly	Thr	Glu	Asp	Ser	
111							195			200					205		
113	ggc	acc	aca	gtg	ctg	ttg	ccc	ctg	gtc	att	ttc	ttt	ggt	ctt	tgc	ctt	672
114	Gly	Thr	Thr	Val	Leu	Leu	Pro	Leu	Val	Ile	Phe	Phe	Gly	Leu	Cys	Leu	
115							210			215					220		
117	tta	tcc	ctc	ctc	ttc	att	ggt	tta	atg	tat	cgc	tac	caa	cg	tgg	aag	720
118	Leu	Ser	Leu	Leu	Phe	Ile	Gly	Leu	Met	Tyr	Arg	Tyr	Gln	Arg	Trp	Lys	
119	225						230			235					240		
121	tcc	aag	ctc	tac	tcc	att	gtt	tgt	ggg	aaa	tcg	aca	cct	gaa	aaa	gag	768
122	Ser	Lys	Leu	Tyr	Ser	Ile	Val	Cys	Gly	Lys	Ser	Thr	Pro	Glu	Lys	Glu	
123							245			250					255		
125	ggg	gag	ctt	gaa	gga	act	act	act	aag	cbc	ctg	gcc	cca	aac	cca	agc	816
126	Gly	Glu	Leu	Gly	Thr	Thr	Thr	Lys	Pro	Leu	Ala	Pro	Asn	Pro	Ser		

RAW SEQUENCE LISTING  
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127	260	265	270		
129	tcc	agt	ccc	act cca ggc ttc acc ccc acc ctg ggc ttc agt ccc gtg	864
130	Phe	Ser	Pro	Thr Pro Gly Phe Thr Pro Thr Leu Gly Phe Ser Pro Val	
131	275		280	285	
133	ccc	agt	tcc	acc ttc acc tcc agc tcc acc tat acc ccc ggt gac tgt	912
134	Pro	Ser	Ser	Thr Phe Thr Ser Ser Ser Thr Tyr Thr Pro Gly Asp Cys	
135	290		295	300	
137	ccc	aac	ttt	gct ccc cgc aga gag gtg gca cca ccc tat cag ggg	960
138	Pro	Asn	Phe	Ala Ala Pro Arg Arg Glu Val Ala Pro Pro Tyr Gln Gly	
139	305		310	315	320
141	gct	gac	ccc	atc ctt gcg aca gcc ctc gcc tcc gac ccc atc ccc aac	1008
142	Ala	Asp	Pro	Ile Leu Ala Thr Ala Leu Ala Ser Asp Pro Ile Pro Asn	
143	325		330	335	
145	ccc	ctt	cag	aag tgg gag gac agc gcc cac aag cca cag agc cta gac	1056
146	Pro	Leu	Gln	Lys Trp Glu Asp Ser Ala His Lys Pro Gln Ser Leu Asp	
147	340		345	350	
149	act	gat	gac	ccc gcg acg ctg tac gcc gtg gtg gag aac gtg ccc ccg	1104
150	Thr	Asp	Asp	Pro Ala Thr Leu Tyr Ala Val Val Glu Asn Val Pro Pro	
151	355		360	365	
153	ttg	cgc	tgg	aag gaa ttc gtg cgg cgc cta ggg ctg agc gac cac gag	1152
154	Leu	Arg	Trp	Lys Glu Phe Val Arg Arg Leu Gly Leu Ser Asp His Glu	
155	370		375	380	
157	atc	gat	cg	ctg gag ctg cag aac ggg cgc tgc ctg cgc gag gcg caa	1200
158	Ile	Asp	Arg	Leu Glu Leu Gln Asn Gly Arg Cys Leu Arg Glu Ala Gln	
159	385		390	395	400
161	tac	agc	atg	ctg gcg acc tgg agg cgg cgc acg ccg cgg cgc gag gcc	1248
162	Tyr	Ser	Met	Leu Ala Thr Trp Arg Arg Arg Thr Pro Arg Arg Glu Ala	
163	405		410	415	
165	acg	ctg	gag	ctg gga cgc gtg ctc cgc gac atg gac ctg ctg ggc	1296
166	Thr	Leu	Glu	Leu Leu Gly Arg Val Leu Arg Asp Met Asp Leu Leu Gly	
167	420		425	430	
169	tgc	ctg	gag	gac atc gag gag gcg ctt tgc ggc ccc gcc gcc ctc ccg	1344
170	Cys	Leu	Glu	Asp Ile Glu Ala Leu Cys Gly Pro Ala Ala Leu Pro	
171	435		440	445	
173	ccc	gcg	ccc	agt ctt ctc aga tga	1368
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175	450		455		
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180	<212>	TYPE:	PRT		
181	<213>	ORGANISM:	Homo sapiens		
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187	Glu	Leu	Leu	Val Gly Ile Tyr Pro Ser Gly Val Ile Gly Leu Val Pro	
188	20		25	30	
190	His	Leu	Gly	Asp Arg Glu Lys Arg Asp Ser Val Cys Pro Gln Gly Lys	
191	35		40	45	
193	Tyr	Ile	His	Pro Gln Asn Asn Ser Ile Cys Cys Thr Lys Cys His Lys	

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194	50	55	60
196	Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly Gln Asp Thr Asp		
197	65	70	75 80
199	Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser Glu Asn His Leu		
200	85	90	95
202	Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu Met Gly Gln Val		
203	100	105	110
205	Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr Val Cys Gly Cys Arg		
206	115	120	125
208	Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu Phe Gln Cys Phe		
209	130	135	140
211	Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu Ser Cys Gln Glu		
212	145	150	155 160
214	Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu		
215	165	170	175
217	Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser Leu Glu Cys Thr		
218	180	185	190
220	Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr Glu Asp Ser		
221	195	200	205
223	Gly Thr Thr Val Leu Leu Pro Leu Val Ile Phe Phe Gly Leu Cys Leu		
224	210	215	220
226	Leu Ser Leu Leu Phe Ile Gly Leu Met Tyr Arg Tyr Gln Arg Trp Lys		
227	225	230	235 240
229	Ser Lys Leu Tyr Ser Ile Val Cys Gly Lys Ser Thr Pro Glu Lys Glu		
230	245	250	255
232	Gly Glu Leu Glu Gly Thr Thr Lys Pro Leu Ala Pro Asn Pro Ser		
233	260	265	270
235	Phe Ser Pro Thr Pro Gly Phe Thr Pro Thr Leu Gly Phe Ser Pro Val		
236	275	280	285
238	Pro Ser Ser Thr Phe Thr Ser Ser Thr Tyr Thr Pro Gly Asp Cys		
239	290	295	300
241	Pro Asn Phe Ala Ala Pro Arg Arg Glu Val Ala Pro Pro Tyr Gln Gly		
242	305	310	315 320
244	Ala Asp Pro Ile Leu Ala Thr Ala Leu Ala Ser Asp Pro Ile Pro Asn		
245	325	330	335
247	Pro Leu Gln Lys Trp Glu Asp Ser Ala His Lys Pro Gln Ser Leu Asp		
248	340	345	350
250	Thr Asp Asp Pro Ala Thr Leu Tyr Ala Val Val Glu Asn Val Pro Pro		
251	355	360	365
253	Leu Arg Trp Lys Glu Phe Val Arg Arg Leu Gly Leu Ser Asp His Glu		
254	370	375	380
256	Ile Asp Arg Leu Glu Leu Gln Asn Gly Arg Cys Leu Arg Glu Ala Gln		
257	385	390	395 400
259	Tyr Ser Met Leu Ala Thr Trp Arg Arg Arg Thr Pro Arg Arg Glu Ala		
260	405	410	415
262	Thr Leu Glu Leu Leu Gly Arg Val Leu Arg Asp Met Asp Leu Leu Gly		
263	420	425	430
265	Cys Leu Glu Asp Ile Glu Glu Ala Leu Cys Gly Pro Ala Ala Leu Pro		
266	435	440	445

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/898,234B

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Output Set: N:\CRF4\11132002\I898234B.raw

268 Pro Ala Pro Ser Leu Leu Arg  
269 450 455  
272 <210> SEQ ID NO: 3  
273 <211> LENGTH: 483  
274 <212> TYPE: DNA  
275 <213> ORGANISM: Homo sapiens  
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278 <221> NAME/KEY: CDS  
279 <222> LOCATION: (1)..(483)  
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283 Asp Ser Val Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser  
284 1 5 10 15  
286 att tgc tgt acc aag tgc cac aaa gga acc tac ttg tac aat gac tgt 96  
287 Ile Cys Cys Thr Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys  
288 20 25 30  
290 cca ggc ccg ggg cag gat acg gac tgc agg gag tgt gag agc ggc tcc 144  
291 Pro Gly Pro Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser  
292 35 40 45  
294 ttc acc gct tca gaa aac cac ctc aga bac tgc ctc agc tgc tcc aaa 192  
295 Phe Thr Ala Ser Glu Asn His Leu Arg His Cys Leu Ser Cys Ser Lys  
296 50 55 60  
298 tgc cga aag gaa atg ggt cag gtg gag atc tct tct tgc aca gtg gac 240  
299 Cys Arg Lys Glu Met Gly Gln Val Glu Ile Ser Ser Cys Thr Val Asp  
300 65 70 75 80  
302 cgg gac acc gtg tgt ggc tgc agg aag aac cag tac cgg cat tat tgg 288  
303 Arg Asp Thr Val Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr Trp  
304 85 90 95  
306 agt gaa aac ctt ttc cag tgc ttc aat tgc agc ctc tgc ctc aat ggg 336  
307 Ser Glu Asn Leu Phe Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn Gly  
308 100 105 110  
310 acc gtg cac ctc tcc tgc cag gag aaa cag aac acc gtg tgc acc tgc 384  
311 Thr Val His Leu Ser Cys Gln Glu Lys Gln Asn Thr Val Cys Thr Cys  
312 115 120 125  
314 cat gca ggt ttc ttt cta aga gaa aac gag tgt gtc tcc tgt agt aac 432  
315 His Ala Gly Phe Phe Leu Arg Glu Asn Glu Cys Val Ser Cys Ser Asn  
316 130 135 140  
318 tgt aag aaa agc ctg gag tgc acg aag ttg tgc cta ccc cag att gag 480  
319 Cys Lys Lys Ser Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu  
320 145 150 155 160  
322 aat  
323 Asn  
326 <210> SEQ ID NO: 4  
327 <211> LENGTH: 161  
328 <212> TYPE: PRT  
329 <213> ORGANISM: Homo sapiens  
331 <400> SEQUENCE: 4  
332 Asp Ser Val Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser  
333 1 5 10 15

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/898,234B

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Input Set : A:\98385Iseq.txt  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:23; N Pos. 344,4157,5135,6255  
Seq#:28; Xaa Pos. 4  
Seq#:29; Xaa Pos. 7  
Seq#:32; Xaa Pos. 1,2  
Seq#:36; Xaa Pos. 6,10,11,12  
Seq#:41; Xaa Pos. 9,10,11  
Seq#:43; Xaa Pos. 6,18  
Seq#:44; Xaa Pos. 16,17  
Seq#:46; Xaa Pos. 7

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23  
Seq#:1; Line(s) 24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43  
Seq#:1; Line(s) 44,45,46,47,48,49,50,51,52,53,54,55,56,57,58,59,60,61,62,63  
Seq#:1; Line(s) 64,65,66,67,68,69,70,71,72,73,74,75,76,77,78,79,80,81,82,83  
Seq#:1; Line(s) 84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99,100,101,102  
Seq#:1; Line(s) 103,104,105,106,107,108,109,110,111,112,113,114,115,116,117  
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Seq#:1; Line(s) 178  
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